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Q gene exon 5

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CR1-L

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R gene exon 2

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U gene exon 1

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U gene exon 2

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U gene exon 3

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U gene exon 4

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U gene exon 5

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U gene exon 6

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71851	AGCTGAAACAA	CTCAGAGTTT	GCATCTGTGA	AACAGCAGCT	GCAATATGGA
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72351	CCTAATTACT	GTTCAGAATA	AAGCACTACA	AAACCTGTGT	CAAATGTCTG
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72651	TGGTTCTTCA	GATCTGAAAG	AACCTAGAGA	TTTCTAAATG	GGAACATACA
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73201	ATCACTGCGG	TGTCCCTG	TTCTCCCATT	GGTCTGGCA	TTGCACGTGG
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73301	TGGTTGCCA	CAGTCTGTGC	AGTTTCTCTG	TGTTTCAGCA	CTGACTCGTG
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CR1 - GG

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CR1 - b

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MAR

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V gene exon 1

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CR1 - c

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V gene exon 2

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V gene exon 6

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Y:OV-1 element

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SDRE fragment

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127851 TAGTTTTCA CAGTCTGTGTC ATGATCTGTC ACAGACCAAG GCAGCACAGC
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 NRE: A, B, C regions
 131401 TCCAAGGTTC CCACATTCTT CTGTTTCTT AAAGATCCC TTATCTGGT
 silencer (common site)
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 Ovalbumin exon L
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 Ovalbumin exon 4
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 135901 GAAGAACAAAG ATTGCAGTTG CAAAAACAAA CCAATGTCCT TGCCCCCTACA

135951 TCCTCTCCC CATAAATTCT ACATTCTCTA TCTACCTTGT GCTTGCCAAC
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Ovalbumin exon 5

136051 AAATGTCCCT CAGCCAAGCT CCGTGGATT TCCTAACTGCA ATGGTTCTGG
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 136201 AGAGATGTAA TCTAGAATTG ATGAAGAGAG TAGACATGTT GTTATATGAA
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Ovalbumin exon 6

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 136601 CCTGGAGCTT CCATTTGCCA GTGGGACAAT GAGCATGTTG GTGCTGTTGC
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 136751 CAGTTATCCT AAAACATTG ACCAGCACAA ATTTCATCAT GATTCAATAT
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Ovalbumin exon 7

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 138451 CAGAGAGCCT GAAGATATCT CAAGCTGTCC ATGCAGCACA TGCAAGAAATC

138501 AATGAAGCAG GCAGAGAGGT GGTAGGGTCA GCAGAGGCTG GAGTGGATGC
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 138601 TCAAGCACAT CGCAACCAAC GCCGTTCTCT TCTTTGGCAG ATGTGTTCC
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W gene exon 6

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MAR-like element

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MENT exon 7

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Z1 gene exon 2

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 184751 TGCTGATGCA AGTGTGTCAAG AGATTACGTG GACTCCACAG AAGGTATTTG
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 185551 TTCTTAATAA CAAAGAGGAA GATAATTCCCT AGCTCAGATG GCTAATAAAA

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 186601 GTCAGAATGC TTTTCTTTCT GCTGTGTTTC TACCACTAAA TTGATAGAAA
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 189551 ATTTAAACTG GAAGAAAATT ATGATCTGAA ACCCATCCTG AGCAACATGG
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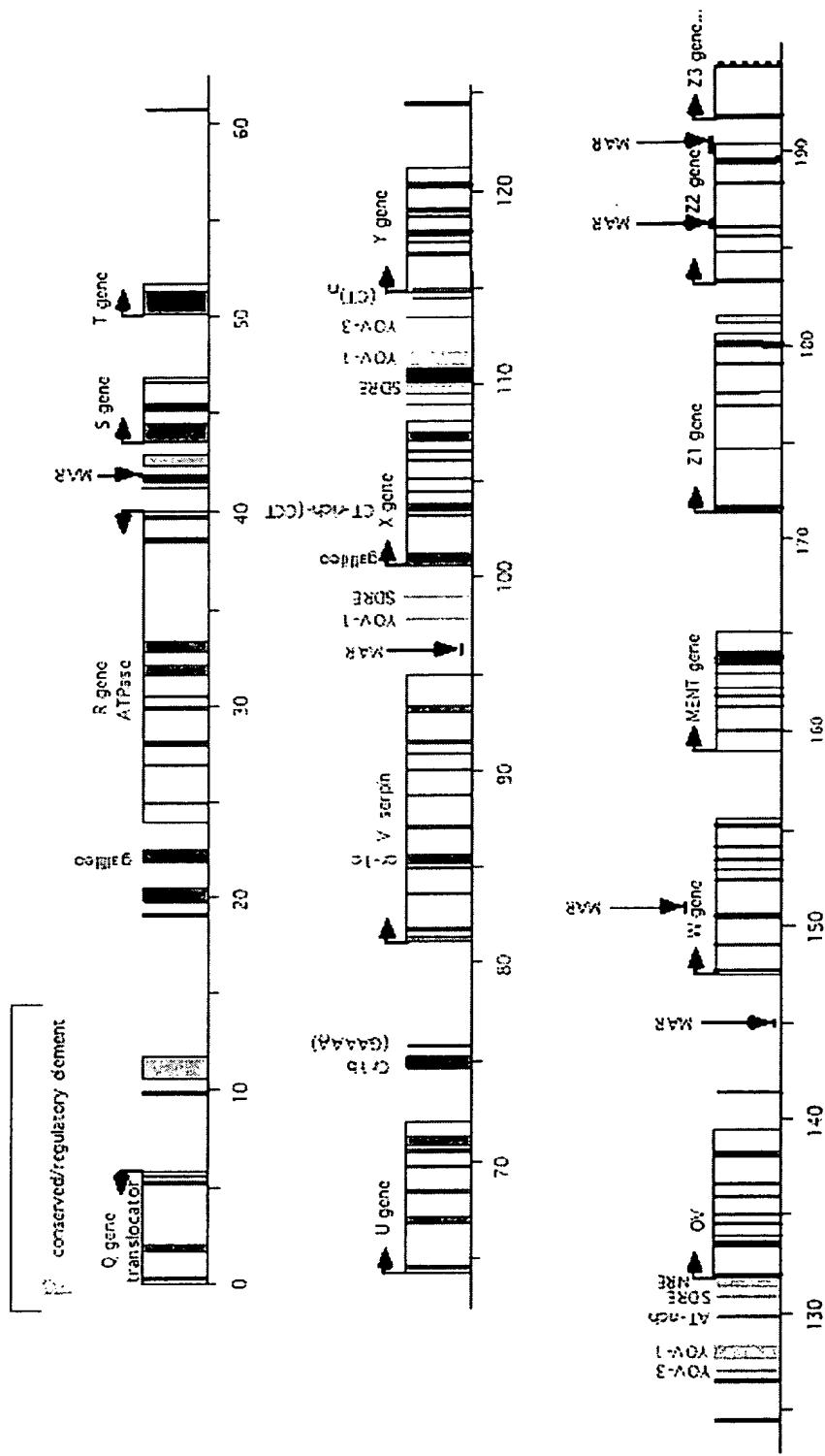
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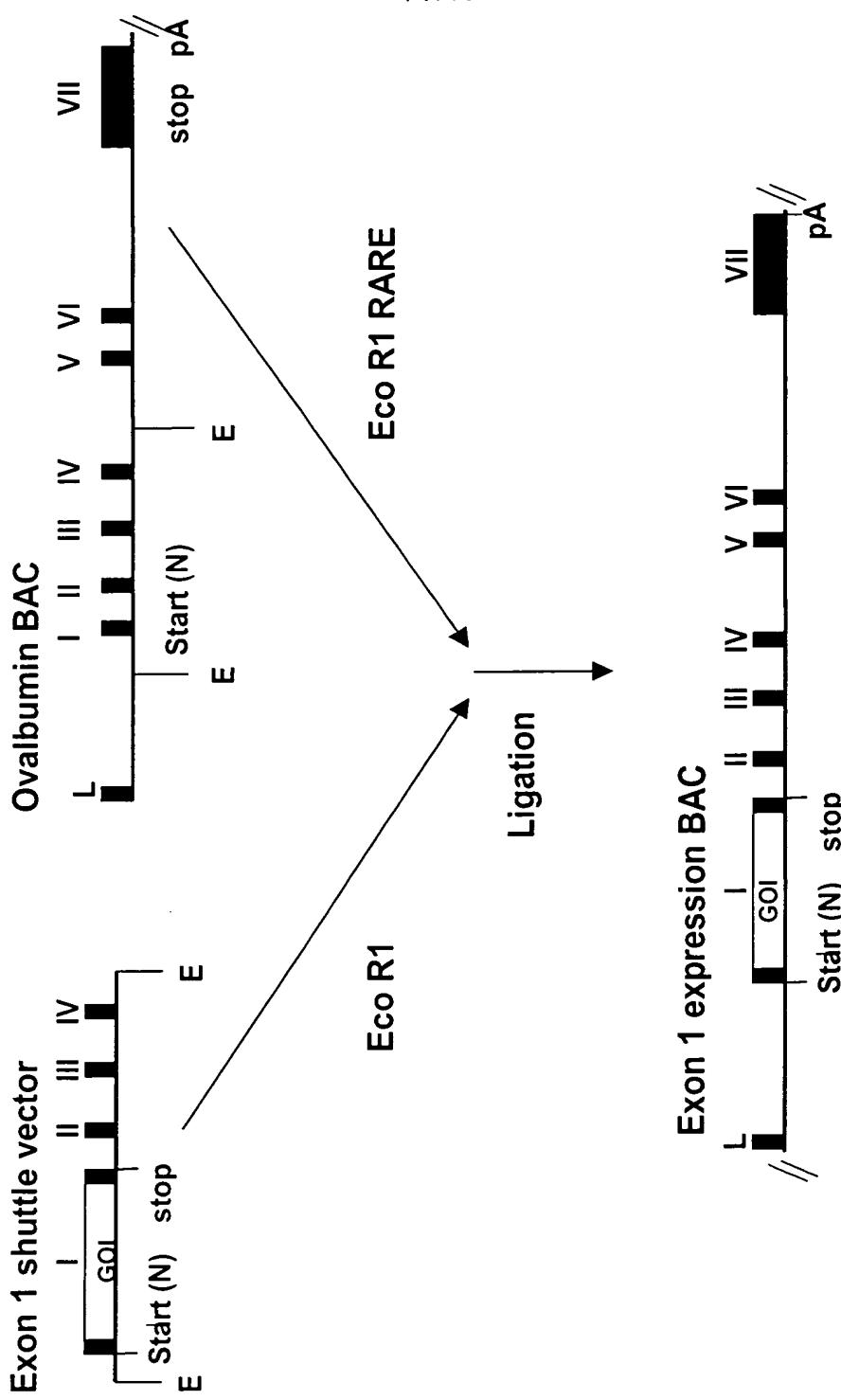
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 192201 *GGTCCTCCTC GGGTCCAGAG GTAATACAGA GACACAGGTG CTGAAGGTCA*
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 192401 *CCAGAGCCCT CCTCTTGTGC TCACTGATAG CCCTTCTTTC TCACTTCATT*
 192451 *TGGGTTAATT GATGAATCTG GAAACTAATT TCACTGATTT ATCAGTCTTA*
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 192651 *AGCTGCCAG GGAAGCTGAG GGTGCCCAT CCCTGGAGGT GTTCAAGACC*
 192701 *AGGTTGGATG GGGCCCTGGG CAGCCTGAGC TGGTGGTGG CTGCCCTGCC*
 192751 *CACAGCAAGG CAGTGGAACT GGGTGGGCTT TAAGTTGAAT TCCAGCCCAA*
 192801 *CCCCATTCTA TGATTCTATG AGCCTTTCC ACAGAGAACT ATTGTTTGC*
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 192951 *CAATACATGG TGATGTTACC CAAGACCCAG TGTTATAGCA GCCAAGCACC*
 193001 *CAGTATTCT GAGGAGCAGA ACTCACGTGT CCATTCTCAT GGTATCCTTA*
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 193151 *GTTGCAAACCT GCAAATTCTC ATGTAGGTAG CAGTGCCAAT TCCCTCTCAG*
 193201 *ACTCATGTTC AAAAGGGACC CTGCCTCTT TTTAATTGCA AAGGCAAACA*
 193251 *CCTGCTAGTG CAAGGGGAAG TATGAAAGAA TTGTCGCTGT AGTTCCATT*
 193301 *AACTTATTG CCCTATGATT AAGTCACCT TTGTATTCCG AACTTTAGGA*

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 195101 TT

Fig. 1

**Fig. 2**

**Fig. 3**

SEQ ID NO: 2

AAAGTCTAGAGTCGGGCGGCCGGCCCTCGAGCAGACATGATAAGATAACATTGATGAG	60
TTGGACAAACCACAACCTAGAATGCAGTGAAAAAAATGCTTATTGTGAAATTGTGAT	120
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ATTCATTTATGTTTCAGGTTCAGGGGAGGTGTGGGAGGTTTAAAGCAAGTAAAAC	240
CTCTACAAATGTGGTAAATCGATAAGGATCCGTCGAGCGGCCGC 285	

FIG. 4**SEQ ID NO: 3**

TGCGATCTGCCTCAGACCCACAGCCTGGCAGCAGGAGGACCTGATGCTGCTGGCTCAG	60
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CAGGGCGTGGCGTGACCGAGACCCCTCTGATGAAGGAGGATAGCATCCTGGCTGTGAGG	360
AAGTACTTCAGAGGATCACCTGTACCTGAAGGAGAAGAAGTACAGCCCCCTGCGCTTGG	420
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Fig. 5

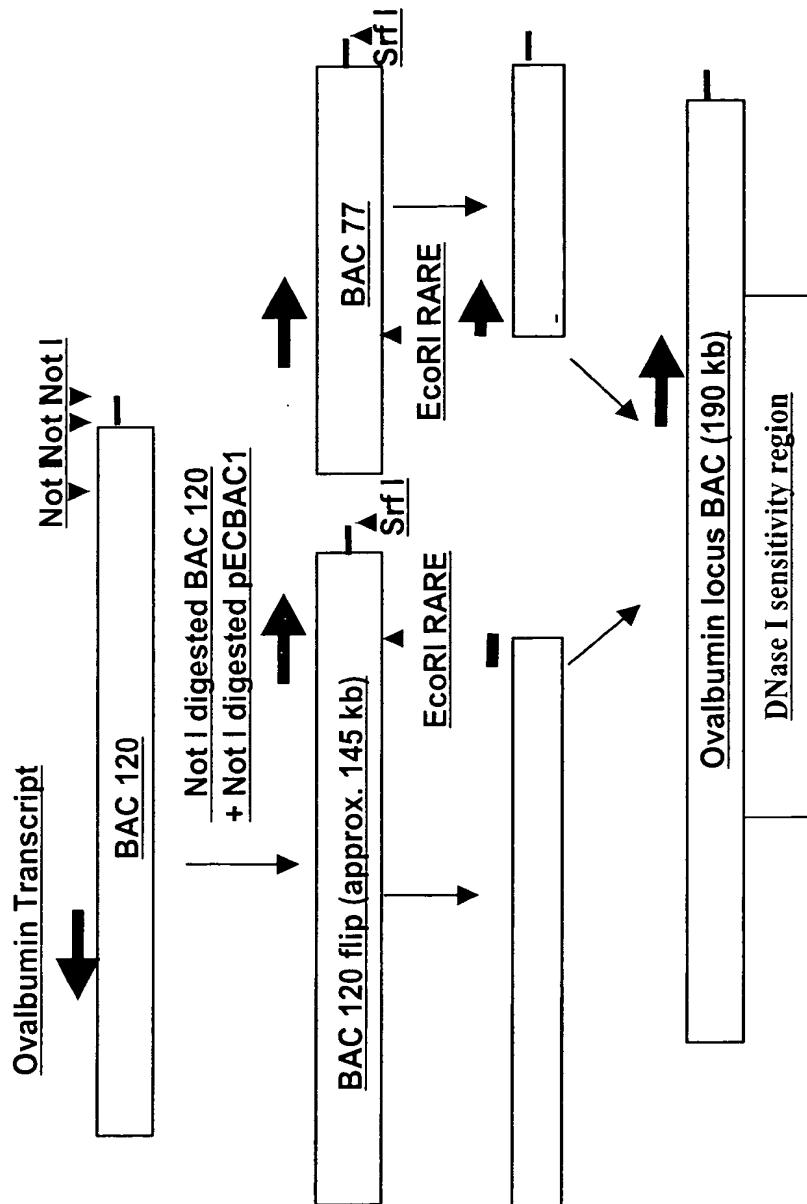


Fig. 6